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Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2007; month=11; day=28; hr=18; min=49; sec=34; ms=540;
]

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Application No: 09776874 Version No: 3.1

Input Set:**Output Set:**

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Finished: 2007-11-28 18:48:19.043
Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 194 ms
Total Warnings: 32
Total Errors: 3
No. of SeqIDs Defined: 47
Actual SeqID Count: 47

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
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W 213	Artificial or Unknown found in <213> in SEQ ID (4)
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W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
E 201	Mandatory field data missing in <223> in SEQ ID (11)
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W 213	Artificial or Unknown found in <213> in SEQ ID (17)
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W 213	Artificial or Unknown found in <213> in SEQ ID (22)
W 213	Artificial or Unknown found in <213> in SEQ ID (23)
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W 213	Artificial or Unknown found in <213> in SEQ ID (26)
W 213	Artificial or Unknown found in <213> in SEQ ID (27)

Input Set:

Output Set:

Started: 2007-11-28 18:48:15.849
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Total Warnings: 32
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Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (28)
W 213	Artificial or Unknown found in <213> in SEQ ID (29) This error has occurred more than 20 times, will not be displayed
E 201	Mandatory field data missing in <223> in SEQ ID (45)

SEQUENCE LISTING

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Vlodavsky , Israel
Feinstein, Elena

<120> POLYNUCLEOTIDE ENCODING A POLYPEPTIDE HAVING HEPARANASE ACTIVITY
AND EXPRESSION OF SAME IN GENETICALLY MODIFIED CELLS

<130> 01/22603

<140> 09776874
<141> 2001-02-06

<150> US 08/922,170
<151> 1997-09-02

<150> US 09/109,386
<151> 1998-07-10

<150> PCT/US98/17954
<151> 1998-08-31

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<211> 1721

<212> DNA

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35 40 45

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Arg Thr Leu Ala Arg Gly Leu Ser Pro Ala Tyr Leu Arg Phe Gly Gly
85 90 95

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100 105 110

Glu Glu Arg Ser Tyr Trp Gln Ser Gln Val Asn Gln Asp Ile Cys Lys
115 120 125

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Val	Ile	Asp	Ser	Val	Thr	Trp	His	His	Tyr	Tyr	Leu	Asn	Gly	Arg	Thr
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Ala	Thr	Arg	Glu	Asp	Phe	Leu	Asn	Pro	Asp	Val	Leu	Asp	Ile	Phe	Ile
305					310					315					320
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Lys	Lys	Val	Trp	Leu	Gly	Glu	Thr	Ser	Ser	Ala	Tyr	Gly	Gly	Gly	Ala
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Pro	Leu	Leu	Ser	Asp	Thr	Phe	Ala	Ala	Gly	Phe	Met	Trp	Leu	Asp	Lys

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ctg ctc ctg ggg ccg ctg ggt ccc ctc tcc cct ggc gcc ctg ccc cga      155
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cct gcg caa gca cag gac gtc gtg gac ctg gac ttc ttc acc cag gag      203
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ccg ctg cac ctg gtg agc ccc tcg ttc ctg tcc gtc acc att gac gcc      251
Pro Leu His Leu Val Ser Pro Ser Phe Leu Ser Val Thr Ile Asp Ala
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aac ctg gcc acg gac ccg cgg ttc ctc atc ctc ctg ggt tct cca aag      299
Asn Leu Ala Thr Asp Pro Arg Phe Leu Ile Leu Leu Gly Ser Pro Lys
           65             70             75

ctt cgt acc ttg gcc aga ggc ttg tct cct gcg tac ctg agg ttt ggt      347
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ggc acc aag aca gac ttc cta att ttc gat ccc aag aag gaa tca acc      395
Gly Thr Lys Thr Asp Phe Leu Ile Phe Asp Pro Lys Lys Glu Ser Thr
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ttt gaa gag aga agt tac tgg caa tct caa gtc aac cag gat att tgc      443
Phe Glu Glu Arg Ser Tyr Trp Gln Ser Gln Val Asn Gln Asp Ile Cys
           115            120            125

aaa tat gga tcc atc cct cct gat gtg gag gag aag tta cgg ttg gaa      491
Lys Tyr Gly Ser Ile Pro Pro Asp Val Glu Glu Lys Leu Arg Leu Glu
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Trp Pro Tyr Gln Glu Gln Leu Leu Leu Arg Glu His Tyr Gln Lys Lys
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ttc aag aac agc acc tac tca aga agc tct gta gat gtg cta tac act      587
Phe Lys Asn Ser Thr Tyr Ser Arg Ser Ser Val Asp Val Leu Tyr Thr
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ttt gca aac tgc tca gga ctg gac ttg atc ttt ggc cta aat gcg tta      635
Phe Ala Asn Cys Ser Gly Leu Asp Leu Ile Phe Gly Leu Asn Ala Leu
           180            185            190

tta aga aca gca gat ttg cag tgg aac agt tct aat gct cag ttg ctc      683
Leu Arg Thr Ala Asp Leu Gln Trp Asn Ser Ser Asn Ala Gln Leu Leu
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Ser Thr Phe Lys Asn Ala Lys Leu Tyr Gly Pro Asp Val Gly Gln Pro	
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Arg Arg Lys Thr Ala Lys Met Leu Lys Ser Phe Leu Lys Ala Gly Gly	
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Thr Ala Thr Arg Glu Asp Phe Leu Asn Pro Asp Val Leu Asp Ile Phe	
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Gly Lys Lys Val Trp Leu Gly Glu Thr Ser Ser Ala Tyr Gly Gly Gly	
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Lys Leu Gly Leu Ser Ala Arg Met Gly Ile Glu Val Val Met Arg Gln	
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